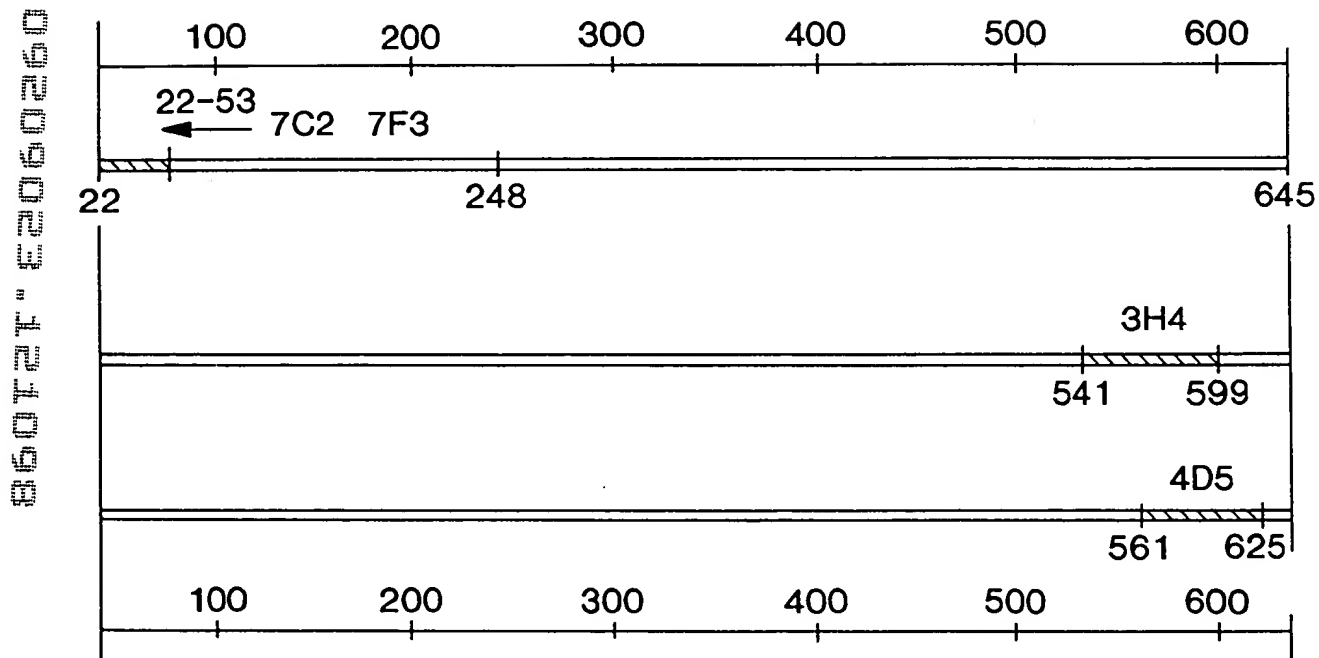


# FIG. 1

3H4	aa 541-599
4D5	aa 529-625
7C2	aa 22-53
7F3	aa 22-53



3H4 epitope (SEQ ID NO:8) 58 residues

VEECRVLQGLPREYVNARHCLPCHPECQPQNGSVTCFGPEADQCVACAHYKDPPFCVAR  
 | |  
 541 599

4D5 epitope (SEQ ID NO:9) (64 residues)

LPCHPECQPQNGSVTCFGPEADQCVACAHYKDPPFCVARCPGSKPDLSYMPIWKFPDEEGACQP  
 | |  
 561 625

860727" 22060260

1 MELAALCRWGLLLALLPPGAASTQVCTGTDMKLRLPA  
38 SPETHLDMLRHLYQGCOVVOGNLELTYLPTNASLSFL  
75 ODIOEVOGYVLIAHNOVROVPLORLRIVRGTOLEFEDN  
112 YALAVLDNGDPLNNTTPVTGASPGGLRELQLRSLTEI  
149 LKGGVLIORNPOLCYODTILWKDIFHKNNQLALTLID  
186 TNRSRA

FIG. 2